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RAW SEQUENCE LISTING DATE: 09/13/2000 PATENT APPLICATION: US/09/649,811 TIME: 18:20:28 Input Set : A:\471cll.app Output Set: N:\CRF3\09132000\1649811.raw 4 <110> APPLICANT: Xu, Jiangchun Lodes, Michael J. Secrist, Heather Benson, Darin R. Meagher, Madeleine Joy King, Gordon E. 12 <120> TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND 13 DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE 16 <130> FILE REFERENCE: 210121.471C11 ENTERED C--> 18 <140> CURRENT APPLICATION NUMBER: US/09/649,811 19 <141> CURRENT FILING DATE: 2000-08-28 21 <160> NUMBER OF SEQ ID NOS: 1083 23 <170> SOFTWARE: FastSEQ for Windows Version 3.0 25 <210> SEQ ID NO: 1 1,000 26 <211> LENGTH: 458 27 <212> TYPE: DNA 28 <213> ORGANISM: Homo sapien 30 <220> FEATURE: 31 <221> NAME/KEY: misc_feature 32 <222> LOCATION: (1)...(458) 33 <223> OTHER INFORMATION: n = A, T, C or G 35 <400> SEQUENCE: 1 W--> 36 ncaggtctgg cggcacctgt gcactcagcc gtcgatacac tggtcgattg ggacagggaa 60 gacgatgtgg ttttcaggga ggcccagaga tttggagaag cggatgaagt tctcctttag 120 ttccgaagtc agctccttgg ttctcccgta gagggtgatc ttgaagtact ccctgttttg 180 agaaactttc ttgaagaaca ccatagcatg ctggttgtag ttggtgctca ccactcggac 240 40 gaggtaactc gttaatccag ggtaactctt aatgttgccc agcgtgaact cgccgggctg 300 41 gcaacctgga acaaaagtcc tgatccagta gtcacacttc.ttttcctaa acaggacgga 360 ggtgacattg tagctcttgt cttctttcag ctcatagatg gtggcataca tcttttgcgg 43 gtctttgtct tctctgagaa ttgcattccc tgccagga 45 <210> SEQ ID NO: 2 46 <211> LENGTH: 423 47 <212> TYPE: DNA 50 <213> ORGANISM: Homo sapien 52 <400> SEQUENCE: 2 cagggtccat aggtgatccg caactctcga gcatttatat acaatagcaa atcatccagt 60 gtgttgtaca gtctataata ctccaacagt ctcccatctg tattcaatgg cgccacccaa 120 tacagtcctt tgtttggatg ctggggagag taatccctac cccaagcacc atatagataa 180 gaaaaccctc tccagttgag ctgaaccaca gacggtttgc tgatgttcac cacaccacca 240 tgaccacage teeetggagt gggaggaggg tggacgacag gggtgttttg atetttagag 300 getteacact ettteagett ggtetteaga geeacgattt eteggegaat ggeaaggaea 360 ttgtttttgt ctagtgtctc aagcttctct accaagagag tcatatttct tatctccacc

60

62 <210> SEQ ID NO: 3 63 <211> LENGTH: 538 64 <212> TYPE: DNA

65 <213> ORGANISM: Homo sapien



RAW SEQUENCE LISTING PATENT APPLICATION: US/09/649,811

DATE: 09/13/2000 TIME: 18:20:28

Input Set : A:\471cll.app
Output Set: N:\CRF3\09132000\1649811.raw

67 (AAA) GEOURNOR 3	
67 <400> SEQUENCE: 3	60
68 ggtctgtcca atggcaacag gaccctcact ctaytcartg tcacaagraa tgayrcagsa 69 msctayraat gtgaaaycca gaacccagtg agtgccarsc gcagtgayyc agtcatcctg	120
69 msctayraat gtgaaaycca gaacccagtg agtgccarsc gcagtgayyc agtcatcctg 70 aatgtcctct atggcccrga tgmccccacc atttcccctc taaacacatm ttaccgwyca	180
71 ggggaaaate tgaacetete etgecaegea geetetaaee cacetgeaca gtactettgg	240
3333	300
	360
	420
	420
75 ctccaacccc gtggaggatg aggatgctgt agccttaacc tgtgaacctg agattcagaa 76 cacaacctac ctgtggtggg taaataatca gagcctcccg gtcagtccca ggctgcag	538
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85 tgcgaacatt gaacggcgtg gattcaatag tgagcttggc agtggtgggc gggttccaga	
86 aggttagaag tgaggctgtg agcaggagcc cctgccaggg gatvcacgca mtctgtgggg	180
87 aggggetgag rggdgweyee atggtetetg etgtetgete tgteeteete tgtggagaag	240
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89 gcccgggcg	309
91 <210> SEQ ID NO: 5	
92 <211> LENGTH: 412	
93 <212> TYPE: DNA	
94 <213> ORGANISM: Homo sapien	
96 <400> SEQUENCE: 5	CO
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98 tatgtatgtg gaatccagaa ctkcagtgag tgcaaaccgc agtgacccag tcaccctgga	120
99 tgtcctctat gggccagaca scccccatca tttccccccc agactcgtct tacctttcgg	180
100 gagcgaacct caacctetee tgccactegg cetetaacce ateceegeag tattettgge 101 kgtateaatg ggatacegea geaacacac caagttetet ttategeeaa aateaegeea	240 300
	360 412
	412
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100 (211) DENGTH: 332 107 (212) TYPE: DNA	
108 <213> ORGANISM: Homo sapien	
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112 ttsaygttkt atgagsysya saatmetgaw getemttyts sakgrwstte kgsatmrgca	120
113 gtsrattcsa catttgggrt akrtymtctc tsgaagysam tgtcakgcag tgrcayccwr	180
114 gkktcwgcwt gcwgtgrgtt amcakcmwtr ywtagkgsgm ayatrattta ramrgtayak	240
115 cymtotomot cytycmocay wtgowcaass mkcacacoto ggoogogaco acgotaagoo 116 cgaattooag cacactggog googttacta gt	300 332
	332
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119 <211> LENGTH: 401 120 <212> TYPE: DNA	
121 <213> ORGANISM: Homo sapien	
123 <400> SEQUENCE: 7	



DATE: 09/13/2000 TIME: 18:20:28 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/649,811

Input Set : A:\471cll.app
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124 125 126 127 128 129	agttgttagt tegggaggtg taagatgagg tggeteettg tetggteeag gatgaegget tgeeetgeea aaegtagage	cctccctggt cccattggga tgataagctg aggtaatgct	agaccaccat cccggatctg atgctgtaat gcttctcgcc	gcgtcccttg gactggttca ttcatcttgg gatgaaggta	aagatggaca ccattgtact ctggcctggc ggtgtaagag	60 120 180 240 300 360
133 134 135	<210> SEQ ID NO: 8 <211> LENGTH: 1151 <212> TYPE: DNA <213> ORGANISM: Homo sa		ctcggtacca	a	·	401
137 138 139						60 120
140 141	ttggtccctc gaggagctcc	agatattaat	ctacctaact	aagtccccag	gtttcttcca	180 240
142	tacaaagaat ccctagacgc	catactgagt	tttaagttcc	ttaattccta	atttaaggct	300
143 144						360 420
144						420
146						540
147	3					600
148	3					660
149						720
150 151						780 840
151						900
153						960
154						1020
155						1080
156						1140
157	ctaccttgga c					1151
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	<211> LENGTH: 604					
	<212> TYPE: DNA					
	<213> ORGANISM: Homo sa <220> FEATURE:	ipien				
	<221> NAME/KEY: misc_fe	ature				
	<222> LOCATION: (1)(
	<223> OTHER INFORMATION		C or G			
169	<400> SEQUENCE: 9					
170	- 3 3 33 3					60
171	2 2					120
172	aattccacat ttgggatagg					180
173 174	totgoatgoa gtgggttaac					240 300
174					-	360
W> 176	gaaattggcg gagagctgcc					420
W> 177	tcctttntta ataacttttg					480
			,,,	,	JJ J J	-

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Output Set: N:\CRF3\09132000\I649811.raw

	178	qaaaacqctq atgcttgttt qaaqatctca agcqcagagt ctqcaagttc atcccctctt	540
	179		600
	180		604
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	189		120
	190		180
	191		240
	192		300
	193		360
	194		420
	195		473
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		<212> TYPE: DNA	
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		<220> FEATURE:	
	203	<221> NAME/KEY: misc_feature	
	204	<222> LOCATION: (1)(411)	
	205	<223> OTHER INFORMATION: n = A,T,C or G	
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	209	cctgatgcag ccacagcage ccgaagggte tcaaaggtgt cctcgatcte aatgatctge	120
	210	tggatgttgt tggtgatggt ggagatgacc ttatcgatga ggtgcaccac cccgttggtt	180
	211	gcatggtggt cggctttyar carccgggca cagttcacag ttacaatccc attaggatag	240
M>	212	tggtggatet nggatgttgg aattetggta catagnaggt gaggggteat geeegtgttt	300
	213	cagotoatoa gtoaggaoto gootgoocao catatggtaa gosgragggo atttgagoag	360
M>	214	ctcaatgttt gacattgctg gaccagggga gttccagcac ttctangang a	411
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		<211> LENGTH: 560	
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		<213> ORGANISM: Homo sapien	
		<400> SEQUENCE: 12	
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	223	33 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3	120
	224	3 33 33 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3	180
	225		240
	226		300
	227		360
	228	3 3 3 - 3 - 3 - 3 - 3 - 3 - 3 - 3 -	420
	229		480
	230 231		540
		caggggaaag tgaaaacaa <210> SEQ ID NO: 13	560
		<211> SEQ 1D NO: 13 <211> LENGTH: 150	
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RAW SEQUENCE LISTING DATE: 09/13/2000 PATENT APPLICATION: US/09/649,811 TIME: 18:20:28

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Output Set: N:\CRF3\09132000\1649811.raw

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235 <212> TYPE: DNA
     236 <213> ORGANISM: Homo sapien
     238 <400> SEQUENCE: 13
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     239
     240 ttgaacacat ttaagatttg agggatataa gggaaaatga tatgaatgtg tatttttact
                                                                                   120
                                                                                   150
     241 caaaataaaa gtaactgttt acgttggtga
     243 <210> SEQ ID NO: 14
     244 <211> LENGTH: 403
     245 <212> TYPE: DNA
     246 <213> ORGANISM: Homo sapien
     248 <400> SEQUENCE: 14
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     250 gctagctatg ggtagccagg tgttacaaag gtgctgctcc ttctccaacc cctacttggt
251 ttccctcacc ccaagcctca tgttcatacc agccagtggg ttcagcagaa cgcatgacac
                                                                                   120
                                                                                   180
                                                                                   240
     252 cttatcacct ccctccttgg gtgagctctg aacaccagct ttggcccctc cacagtaagg
     253 ctgctacatc aggggcaacc ctggctctat cattttcctt ttttgccaaa aggaccagta
                                                                                   300
     254 gcataggtga gccctgagca ctaaaaggag gggtccctga agctttccca ctatagtgtg
                                                                                   360
     255
          gagttctgtc cctgaggtgg gtacagcagc cttggttcct ctg
                                                                                   403
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     258 <211> LENGTH: 688
     259 <212> TYPE: DNA
     260 <213> ORGANISM: Homo sapien
     262 <220> FEATURE:
     263 <221> NAME/KEY: misc_feature
     264 <222> LOCATION: (1)...(688)
     265 <223> OTHER INFORMATION: n = A, T, C or G
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     269
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                                                                                   120
     270 caaaagcaca gaagcacatc acatacacca gcaaggtttc caactactgc actgattaac
                                                                                   180
          tagatactet caatagettt tetatagete gteetagaaa aaaaaattaa atttteattt
                                                                                   240
     271
     272 tottacaagt tocaggotta aacaaaggoa aaaattacat goaacaactg atacactcat
                                                                                   300
          aagttgcaca tatgctccaa ggtctttatt agataacaat aaatgctagc actttgtcac
                                                                                   360
     273
     274 tgccatcaga ttttccttat agtcttagag tcatgtaaat aaaagttcca taatgaaatt
                                                                                   420
     275 aaagaaaatt aatttttcta atcttagatc agttccatag aaaactatta attttttaa
                                                                                   480
     276 agtaggcagt agaaggggt tggtggggg tggaattggt tagtaagtct ggttctaatc
                                                                                   540
                                                                                   600
     277 ttctgagctg cctttggaag gaagttatga ggtagaagat tctactgact tttagtaagg
W--> 278 tggacaatga gagaaaagaa aaagcaggtg cctcatcnnc agatccttnt ggtatttatn
                                                                                  660
W--> 279 tgccangtnc nanntaatnc atanaaag
                                                                                   688
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     282 <211> LENGTH: 408
     283 <212> TYPE: DNA
     284 <213> ORGANISM: Homo sapien
     286 <400> SEQUENCE: 16
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                                                                                  120
     288 agtttatgat tacaggtgga aatctacaag acagtaaaga tgcactgcat ttggcacaaa
     289 caaatggtat gtttttcagt acagttggat gtcgtcctac aagatgtggt gaatttgaaa
                                                                                  180
     290
                                                                                  240
         agaataaccc tgatctttac ttaaaggagt tgctaaatct tgctgaaaac aataaaggga
     291 aagttgtggc aataggagaa tgcggacttg attttgaccc gactgcagtt ttgtcccaaa
                                                                                  300
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Input Set : A:\471cll.app

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L:18 M:270 C: Current Application Number differs, Wrong Format L:36 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 L:212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 L:214 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 L:278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 L:279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 L:365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 L:384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 L:412 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 L:430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 L:431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 L:433 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 L:434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 L:451 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:26 L:452 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 L:454 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 L:467 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:468 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:469 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:470 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:471 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:472 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:473 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 L:489 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 L:590 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 L:591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 L:592 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 L:593 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 L:620 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 L:621 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 L:622 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 L:623 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 L:624 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 L:625 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 L:626 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:37 L:653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 L:656 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 L:688 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 L:691 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:41 L:692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 L:749 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 L:766 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46 L:768 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46 L:813 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 L:826 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50

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Input Set : A:\471cll.app
Output Set: N:\CRF3\09132000\1649811.raw

L:827 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 L:828 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 L:829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50